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GenCore version 5.1.3

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OM nucleic - nucleic search, using sw model

Run on: December 13, 2002, 15:41:52; Search time 268 Seconds

(without alignments)

1378.089 Million cell updates/sec

1 US-09-659-737A-1

Perfect score: 164
Sequence: 164
Sequence: 1 gcaccgggacatcaaggcag.....tgcctggatggccccagaag 164
Scoring table: IDENTITY_NUC
Gapop 10.0, Gapext 1.0

Searched: 2185239 seqs, 112599159 residues
Total number of hits satisfying chosen parameters: 4370478
Minimum DB seq length: 2000000000

Maximum Match 100%
Listing first 45 summaries

Database: N_Geneseq_101002:*
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N\_Geneseq\_101002:\*

| N\_Geneseq\_101002:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1980.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1981.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1982.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1983.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1989.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1989.DAT:\*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA1999.DAT:\*
| SIDS2/gcgdata/

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Human MLK4 partial cDNA encoding huma Novel human protei Novel human protei Human kinase (PKIN Human PKIN-12 CDNA Human polynucleoti Human cDNA differe
SUMMARIES	AARFOU487 AARFG13 ABN86357 ABN86358 AAD18824 AAD18824 AAD345458 AAD343874
DB	22 22 24 44 22 22 22 24 25 25 25 25 25 25 25 25 25 25 25 25 25
% Query Match Length DB ID	164 3066 3111 3111 3518 3518 3538 444 3558
% Query Match	100.0 933.2 933.2 933.2 57.4 57.4 533.0
Score	152.8 152.8 152.8 152.8 152.8 94.2 87
Result No.	1100410180

8 13	nase kinase; UNKK; skin damaye; inflammation; atory; antipsoriatic;
AAD36139 AAB17018 AAB17018 AAB17018 AAAC49755 AAAC49755 AAAC49751 AAT03096 AAT03096 AAT03094 AAT03094 AAT03034 AAT03034 AAT03328 AAT99250 AAT99249 AAAC43328 AAAC433328 AAAAC433328 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	l kinase kinase k radiation; skin d antiinflammatory; lifiers
447044040404040404040404040404040404040	try) (A. A. Con/Qua con/Qua al al S5029.
33558 3454 404 404 404 404 404 404 137 137 137 1008 3427 1008 10	(first entry)  rtial cDNA.  rtial cDNA.  c. Jun N-terminal kinas  e; ultraviolet radiati dioprotective; antiinf  Location/Qualifiers 2.163 /*tag- a /partial  2000EP-0307866.  99US-0155029.  NEW YORK STATE.
4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	(first partial of the condition of the c
6.77 7.72 8.72 7.72 8.72 7.72 8.72 7.72 8.72 7.72 8.72 7.72 8.72 7.72 8.72 7.72 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73 8.73	30487; 30487; MAY-20 an Mik tein k riasis nerary D sapi D Sapi MAR-20 SEP-19
1386 11111111111111111111111111111111111	AAF. 29-1 Humm MLK PLOW PLOY PLOY PLOY PLOY PLOY PLOY PLOY PLOY
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02-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein Kinase, enzyme; cytostatic; nootropic; neuroprotective; human; antiparkinsonian; virucide; antibacterial; antifungal; antimigraine; analygesic; hypotensive; hypertensive; immunosuppressive; antialergic; antipheumatic; antiarthritic; ophthalmological; anorectic; osteopathic; thrombolytic; antiarteriosclerotic; antiasthmatic; vasotropic; antidiabetic; gene therapy; ss.
                                                                                                                                                                                                                    The present sequence is that of a partial CDNA encoding MLK4 (see AAB20333), lof 4 novel c-Jun N-terminal kinase kinases consists of the invention. CDNAs encoding JNKKKS were isolated from human foreskin epidermal keratinocyte cDNA by PCR amplification using primers (see AAB30494-97) based on regions of homology between the kinase domains of JNKKS or JNKKKS. MLK4 was 50-78% identical to members of the MLK family of JNKKKS. MLK4 was 50-78% identical to members of the MLK family of JNKKKS. The MLK4 gene is cxpressed in keratinocytes, kidney and pancress, but not in brain, placenta, lung, liver or skeletal muscle. The transcript size is products are useful for elucidation of the components involved in the cellular response to ultraviolet radiation. They can be used in drug discovery, by screening for compounds that affect the cellular JNKKK. Particularly useful are drugs that reduce UV and the component and procession of a gene encoding a JNKKK. Particularly useful are drugs that reduce UV and the component of the spread of the skin, inflammation and psoriasis, and
                                                                                                     New polynuclectides encoding c-Jun N-terminal kinase kinases kinases in e. MIKM, PAK4, associated with skin damage for use in drug screening and development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 22; Length 164;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121 CAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 164 BP; 55 A; 33 C; 44 G; 32 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 100.0%; Score 164; DB 22; Best Local Similarity 100.0%; Pred. No. 1.9e-45; Matches 164; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA encoding human protein kinase SGK067.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              drugs that enhance wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH46913 standard; cDNA; 2157 BP.
                                                                                                                                                                                   Claim 3; Page 22; 51pp; English.
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Blumenberg M, Gazel AM;
                                     WPI; 2001-236883/25.
                                                             P-PSDB; AAB20333
                                                                                                                                             and development
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200155356-A2
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The invention provides human protein kinases and protein kinase-like enzymes and polynuclectides encoding the polypeptides. The kinase polypeptides and polynuclectides encoding the polypeptides. The kinase polypeptides and their modulators are useful for treating a disease or disorder such as cancer, immune-related diseases, cardiovascular disease, brain or neuronal associated disease and metabolic disorders, including cancers of tissues, cancers of the metapoletic origin, diseases of the central nervous system, diseases of the peripheral nervous system, and indicated of disease, parkinson's disease, multiple sclerosis, amyotrophic lateral sclerosis, viral infections, infections quant, sexual dysfunction, bacteria and fungi, ocular diseases, migraines, pain, sexual dysfunction, hypertension, psychotic disorders, neurological disorders, dyskinesias, metabolic disorders, autoimmunity, atherosclerosis, psoriasis, osteoarthritis, asthma, chronic inflammatory pelvic diseases rheumatoid arthritis, metabolic disorders cuch as diabetes, obesity, cardiovascular diseases such as reperfusion injury, coronary thrombosis, clotting disorders and atherosclerosis, chronic inflammatory boyelerosis, cancertion, coular diseases such as glaucom, retinopathy and macular degeneration, psychiatric and metulological disorders such as anxiety, schiatophrenia, provinitaring and metulogical disorders such as anxiety services.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dementia, manic depression, etc. The polynucleotides are useful in gene therapy techniques to treat the above mentioned disorders. Sequences AAH46891-46922 represent human protein kinases encoding cDNA molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kinase polypeptides useful for treating cancers, Alzheimer's disease, viral infections, diabetes, obesity, organ transplant rejection and rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plowman G, Whyte D, Manning G, Sudarsanam S, Martinez R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2157 BP; 514 A; 611 C; 638 G; 394 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  900 CAAAATGAGCACAGGAGCACCTATGCCTGGATGGCCCCCGAAG 943
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                                                                                                                                                          2000US-0183173.
2000US-0190162.
2000US-0193404.
25-JAN-2001; 2001WO-US02337.
                                                                                    2000US-0178078
                                                                                                                                                                                                                                                                                                          2000US-0247013
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                                                                                                                                                                                                                                                                                                                                                                                               (SUGE-) SUGEN INC.
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                                                                                                                  31-JAN-2000;
17-FEB-2000;
17-MAR-2000;
29-MAR-2000;
                                                                                    25-JAN-2000;
                                                                                                                                                                                                                                                                                                     13-NOV-2000;
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The invention relates to an isolated polypeptide with signal sequences which allow it to be secreted extracellularly or membrane associated. The activity of polypeptides of the invention may be described as, cytostatic, vulnerary, antiarteriosclerotic, antiparkinsonian, mostropic, neuroprotective, antialoer, virucide, antiparkinsonian, mostropic, cardiant, antiuloer, virucide, antiphyroid, cerebroprotective, anorectic, and metabolic Polypeptides and polynucleotides of the invention are useful in the treatment, or as a vaccine in the prevention of, cancer, wound healing disorders, infection, atherosclerosis, Parkinson's disease and Alzheimer's disease, antoimmune disorder, haematopoietic disorder, inflammation, neoplastic diseases, nervous system related disorders and cardiovascular disorders, pancreatitis, respiratory disorders and edvelopmental abnormality, gastroinestian ulceration, neuropathy, haematological diseases, metabolic diseases, sperm dysfunction, thyroid disorders e.g., hypothyroidism, brain damages, scritis, come photo-claranduction deficiency, neurological diseases, stroke, anajogenesis, ovultation disorders, diseases in the spinal cord, thyroid gland, heart,
                                                                                                                                                                                  Human; cytostatic; vulnerary; antiarteriosclerotic; antiparkinsonian;
Mootropic; neuroprotective; immunosuppressive; haemostatic;
Mootropic; neuroprotective; immunosuppressive; haemostatic;
Malinflammatory; cardiant; antiulcer; virucide; antithyroid;
cerebroprotective, anorectic; metabolic; vaccine; cancer; infection;
wound healing disorders; atherosclerosis; Parkinson's disease;
Malcheimer's disease; autoimmune disorder; haematopoietic disorder;
Inflammation; neoplastic disease; nervous system disorder;
Alcheimer disorders; pancreatitis; respiratory disorder;
Mardiovascular disorders; pancreatitis; respiratory disorder;
Macholovascular disorders; pancreatitis; respiratory disorder;
Macholovascular disorders; pancreatitis; respiratory disorder;
Macholovascular disorders; pancrease; sperm dysfunction;
hyperproliferation; systemic autobilit disease; hyper immunity;
developmental abnormality; gastrointestinal ulceration; neuropathy;
hyperproliferation deficiency; neurological disease; stroke;
maylogenesis; ovulation disorder; spinal cord; thyroid gland; heart;
maylogenesis; ovulation disorder; spinal cord; thyroid gland; heart;
maylogenesis; hymph node; muscular system; obesity; anorexia;
maylogenesis; pamph node; muscular system; obesity; anorexia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cogswell JP, Kabnick KF, L
Smith RF, Strum JC, Xie Q;
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30-JAN-2001; 2001US-264922P.
06-FEB-2001; 2001US-266797P.
19-MAR-2001; 2001US-27698PP.
                                                                                                       Novel human gene. SEQ ID 36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-DEC-2001; 2001WO-US49232.
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10-SEP-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GLAX ) GLAXO GROUP LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder -
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trachea, thymus, lymph node and muscular system, obesity, anorexia, growth abnormalities, and alleviation of precocious puberty. The sequences given in records ABQ86130-ABQ86184 represent novel human cDNA's of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel human protein (NKP), kinase that shares structural similarity with animal kinases. The kinase polynucleotides are useful in therapeutic, diagnostic and pharmacogenomic applications and for identifying compounds that modulate, i.e. act as agonists or antagonists of the agene expression or gene product activity. The present sequence represents the NHP kinase coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel human kinase polynucleotide encoding a protein that shares structural similarity with animal kinases for therapeutic, diagnostic
                                                                                                                                                                                                            61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGAGACCAC 120
                                                                                                                                                                                                                                                                                  1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGACAT 60
                                                                                                                        93.2%; Score 152.8; DB 24; Length 3066; 95.7%; Pred. No. 4e-41; 1ive 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/transl_except= (pos: 2773..2775, aa: Xaa)
/product= "NHP kinase"
                                                                                         Sequence 3066 BP; 738 A; 881 C; 852 G; 595 T; 0 other;
                                                                                                                                                                                                                                                                                                                                  121 CAAAATGAGCACAGGAGGACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                      /note= "Xaa can be any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human protein; NHP; kinase; human; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human protein (NHP) kinase coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 36-37; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and pharmacogenomic applications -
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABN86357 standard; DNA; 3111 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (LEXI-) LEXICON GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-DEC-2001; 2001WO-US47606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-DEC-2000; 2000US-254744P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hu Y, Kieke JA, Donoho G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-OCT-2002 (first entry)
                                                                                                                                                             Matches 157; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-566739/60.
P-PSDB; ABB80923.
                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200255685-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN86357;
                                                                                                                              Query Match
                                                                                                                                              Best Local
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                                                                                                                                                                                                                                                                            61 CIGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                                                                                                            Novel human kinase polynucleotide encoding a protein that shares structural similarity with animal kinases for therapeutic, diagnostic
                                                                                                                                                                                             0; Gaps
                                                                                                                                                                 1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACAT
                                               93.2%; Score 152.8; DB 24; Length 3111; ilarity 95.7%; Pred. No. 4.1e-41; Conservative 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93.2%; Score 152.8; DB 24; Length 3518; 95.7%; Pred. No. 4.3e-41; trive 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3518 BP; 811 A; 1043 C; 976 G; 680 T; 8 other;
Sequence 3111 BP; 740 A; 905 C; 864 G; 594 T; 8 other;
                                                                                                                                                                                                                                                                                                                                                                                                                     121 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel human protein; NHP; kinase; human; gene; ds.
                                                                     4.1e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence represents the NHP kinase encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human protein (NHP) kinase encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 39-41; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
264..3374
/*tag = a
/product="kinase"
/note="NHP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and pharmacogenomic applications -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN86358 standard; DNA; 3518 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LEXI-) LEXICON GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-DEC-2001; 2001WO-US47606.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hu Y, Kieke JA, Donoho G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-566739/60.
                                                                        Best Local Similarity
Matches 157; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB; ABB80923.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABN86358;
                                                     Query Match
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Gaps

Best Local Similarity 95.7 Matches 157; Conservative

Similarity

Query Match

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Human kinase; PKIN; gene therapy; adenocarcinoma; immune disorder; gout; cancer; allergy; sarcoma; leukaemia; acquired immune deficiency syndrome; AIDS; Addison's disease; mircobial infection; inflammation; osteoporosis; atherosclerosis; cardiovascular disease; myocardial infarction; anaemia; myasthenia gravis; cirrhosis; cartaract; growth and development disorder; selzure disorder; pulmonary embolism; Gaucher's disease; lipid disorder; lipid storage disease; pick's disease; Tay-sachs disease; renal disease; obesity; restorative therapy; immunomodulatory; vaccine; cardiovascular; antimicrobial; oytostatic; antilifiammatory; asthma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to human kinases (PKIN) and the nucleic acids encoding them. PKIN is used as vaccine and in gene therapy. PKIN is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yue H, Gandhi AR, Tribouley CM, Kearney L, Griffin JA, Nguyen DB; Bandman O, Lu DAM, Lal P, Burford N, Khan FA, Walia NK, Yao MG; Patterson C, Burrill JD, Marcus GA, Zingler KA, Recipon SA, Lu Y; Policky JL, Thornton M, Tang YT, Hafalia A, Elliott VS, Baughn MK; Walsh RT, Ramkumar J, Borowsky ML, Au-young J, Hillman JL;
                   61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human kinases and nucleic acids, useful for preventing diagnosing and treating cancers, inflammation and immune disorders -
1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACAT 60
                                                                                                                                                         121 CAAAATGAGCACAGGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /product= "Human PKIN-9 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 5; Page 157-158; 166pp; English.
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1..3141
                                                                                                                                                                                                                                                               AAD18824 standard; cDNA; 3141 BP.
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28-APR-2000; 2000US-200226P.
05-MAY-2000; 2000US-20339P.
11-MAY-2000; 2000US-203505P.
18-MAY-2000; 2000US-205564P.
26-MAY-2000; 2000US-20739P.
01-JUN-2000; 2000US-208795P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-APR-2001; 2001WO-US12992.
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                                                                                                                                                                                                                                                                                                                                                                       Human kinase (PKIN)-9 cDNA.
                                                                                                                                                                                                                                                                                                                                     (first entry)
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Gururajan R;
                                                                                                                                                                                                                                                                                                                                     18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-2001.
                                                                                                                                                                                                                                                                                                   AAD18824;
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adenocarchioma, lenkaemia, sarrooma, immune disorder, Addison's disease, adenocarchioma, lenkaemia, sarrooma, immune disorder, Addison's disease, acquired immune deficiency syndrome (AIDS), anaemia, asthma, allerqies, cquit, microbial infections, cariotowascular disease and/or inflammation, myasthenia gravis, atherosclerosis, cirrhosis, osteoporosis, myocardial infarction, cateract, growth and development disorder, setzure disorder, pulmonary embolism, Gaucher's disease, ripid disorder, ilpid storage disease, pick's disease, pray-Sachs disease, renal disease and obesity.

Expression by rectifying mutations or deletions in a patient's genome that affect the activity of PXIN by expressing inactive proteins or to supplement the patients own production of PXIN nucleic acids may be used to produce the PXIN polypeptide, by inserting the nucleic acids into a host cell and culturing the cell to express the protein. PXIN nucleic acid and its complementers sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar in need of restorative therapy. The present sequence is human PXIN-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acquired immune deficiency syndrome; thymic hypoplasia; Crohn's disease; asthma; neurological disorder; epilepsy; Charcot-Marie-Tooth disease; AIDS; selzures; cell proliferative disorder; cancer; adenocarcinoma; leukaemia; lymphoma; melanoma; myeloma; sarcoma; developmental disorder; Down's syndrome; gene therapy; protein therapy; cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; kinase; enzyme; PKIN-12 protein; immune system disorder; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 CACCGGGACATCAAGGCAGGAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC
used in the prevention, diagnosis and treatment of diseases cancers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 57.4%; Score 94.2; DB 22; Length 3141; Best Local Similarity 73.6%; Pred. No. 2.3e-21; Matches 120; Conservative 0; Mismatches 43; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /product= "Human mature PKIN-12 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3141 BP; 685 A; 941 C; 942 G; 573 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /product= "Human PKIN-12 protein"
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52..3291
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AAD34309
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; hematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                 New human kinases, useful for diagnosing, treating or preventing immune system disorders (e.g. Crohn's disease), neurological disorders (e.g. epilepsy), or cell proliferative disorders (e.g. cancers such as leukemia or lymphoma)
                                                                                                                                                                                                                               Tribouley CM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                                                                                                                             Gandhi AR;
                                                                                                                                                                                                           Nguyen DB, Walia NK, Hafalia AJA, Yao MG, Gandhi AF

7. Ding L, Patterson C, Yue H, Baughn MR, Tribouley

Elliott VS, Lu Y, Ison CH, Au-Young J, Tang YT;

Burrill JD, Marcus GA, Zingler KA, Lu DAM, Lal PG;

Warren BA, Kearney L, Policky JL, Thangavelu K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   57.4%; Score 94.2; DB 24; Length 3538; 73.6%; Pred. No. 2.4e-21; tive 0; Mismatches 43; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3538 BP; 763 A; 1055 C; 1062 G; 658 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122 AAAATGAGCACAGGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 91; Page 207-208; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human polynucleotide SEQ ID NO 5518.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAI85458 standard; cDNA; 444 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               encoding human PKIN-12 protein.
             20000S-229873P.
20000S-231357P.
20000S-232654P.
20000S-234902P.
20000S-23699P.
                                                                                                                                                                         (INCY-) INCYTE GENOMICS INC.
                                                                                                                                     13-OCT-2000; 2000US-240542P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 73.6%
Matches 120; Conservative
                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-329769/36.
P-PSDB; AAE21717.
                                                                                                                                                                                                                               Gururajan R,
Thornton M, E
Azimzai Y, Bu
Ramkumar J, W
             31-AUG-2000; 2
08-SEP-2000; 2
14-SEP-2000; 2
                                                                        22-SEP-2000;
29-SEP-2000;
06-OCT-2000;
                                                                                                                                                                                                                 Bandman O,
                                                                                                                                                                                                                                                                                                                Burford N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA185458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 8
AA1854584
TO AA18541
XX AA18541
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XX WW HUMEN 1
XX W HUMEN 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO and fifth wipo.int/pub/published_pot_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  74 TIGAAGATTACAGATTITGGGTTGGCGAGGGAATGGCACAGGACCACCAAAATGAGCACA 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              443 AAGTCCAGCAACATATTGATCCTCCAGAAGGTGGAGAATGGAGACCTGAGCAACAAGATT 384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 AAGGCAGGAAATATTTGCTACTTGAGAAGATAGAACATGATGACATCTGCAATAAAACT 73
                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human cDNA differentially expressed in granulocytic cells #445.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, ss, granulocytic cell; DNA chip; bacterial infection; viral infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis; rheumatoid arthritis; glomerulonephritis; asthma; thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 5518; 1399pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       53.0%; Score 87; DB 22; Length 444; 73.5%; Pred. No. 2.7e-19; Live 0; Mismatches 40; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 444 BP; 110 A; 119 C; 111 G; 101 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   323 GCAGGGACGTATGCTTGGATGGCACCCGAAG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       134 GCAGGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK83874 standard; cDNA; 3558 BP.
                                                                                                                                                                                                                                                                                                           Tang YT, Liu C, Drmanac RT;
                                                                                                                                                 26-FEB-2001; 2001WO-US04927.
                                                                                                                                                                                            28-FEB-2000; 2000US-0515126
18-MAY-2000; 2000US-0577409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                WPI; 2001-514838/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                             HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                          P-PSDB; AAO05527
                                                            WO200164835-A2.
                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation
                                                                                                         07-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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The invention relates to detecting (MI) granulocyte (GC) activation (GCA), by detecting the level of expression of gene(s) (Gs) identified by the Analysis as given in the specification, and comparing the expression level in an unactivated of GCA, where differential expression level in an unactivated of GCA, where differential expression of GS is indicative of GCA, agent CGC, where differential expression of GS is indicative of GCA agent alters the expression of at least one gene in GS; (2) screening (M3) of Toran agent capable of modulating (M3) GA by contacting GC with an agent capable of modulating cesponse in a subject, exposure of a subject to a pathogen or sterile inflammation dispection of a there is a least one gene in GS; (2) screening (M3) of Toran agent capable of modulating response in a subject, exposure of a subject to a pathogen or sterile inflammation; dispecting (M4) an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of the gene is indicative of inflammation; or sterile inflammatory disease, by detecting (M5) an inflammatory disease, by contacting a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease, by contacting a tissue having in the tissue. M1 is useful for detecting a tissue having an inflammatory disease, by contacting a tissue having contacting (M5) an inflammation in a tissue, M4 is useful for fererably in an inflammation in a tissue, M4 is useful for response in a subject, exposure of a subject to a pathogen or sterile contammation in a tissue, M4 is useful for response in a subject, exposure of a subject to a pathogen or sterile contammatory disease, crohn's disease, crohn's disease, crohn's disease, crohn's disease, crohn's disease, or of the printed specification, protozoal infection, viral infection or periodontal disease, also bacterial infection, viral infection, protozoal infection, viral infection or treatting one of the above conditions. The present confirmat directly port of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting granulocyte activation by detecting differential expression of genes associated with granulocyte activation, which serves as diagnostic markers that is useful for monitoring disease states and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
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cardiac reperfusion injury; renal reperfusion injury; ARDS; adult respiratory distress syndrome; inflammatory bowel disease; Crohn's disease; ulcerative colitis; periodontal disease; granulocyte activation; chronic inflammation; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 3558;
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69.3%; Pred. No. 1.4e-17;
Live 0; Mismatches 50; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3558 BP; 635 Å; 1211 C; 1161 G; 551 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                 Beazer-Barclay Y, Weissman SM, Yamaga S, Vockley J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID No 445; 114pp; English.
                                                                                                                                                                                                                                                                   03-OCT-2001; 2001WO-US30821.
                                                                                                                                                                                                                                                                                                                 03-OCT-2000; 2000US-237189P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 69.39
Matches 113; Conservative
                                                                                                                                                                                                                                                                                                                                                                   (GENE-) GENE LOGIC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-435328/46.
                                                                                                                                                                      WO200228999-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  drug toxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid; stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous; cytostatic; gene therapy; antineoplastic; Wllm's tumour; addenocarcinoma;
                                                                                                                                                                              62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                        Gaps
                                                                                                                                                   2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 51
                                                                                                        ö
                                                             Length 3558;
                                                        Ouery Watch 50.6%; Score 83; DB 24; Length 35; Best Local Similarity 69.3%; Pred, No. 1.4e-17; Matches 113; Conservative 0; Mismatches 50; Indels
                Sequence 3558 BP; 635 A; 1211 C; 1161 G; 551 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                            1316 CAAATGAGTGCCGGGGCACCTACGCTGGATGCTCCTGAGG 1358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pancreas cancer related gene sequence SEQ ID NO:8355.
                                                                                                                                                                                                                                                                                                                                 122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL70018 standard; DNA; 3454 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAY-2001; 2001WO-US10838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-234009P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAY-2002 (first entry)
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28-SEP-2000;
29-SEP-2000;
29-SEP-2000;
02-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL70018;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method for screening biologically active agent that modulates cancer associated protein kinase function. The invention also relates to a method for diagnosing cancer comprising determining the upregulation of expression of a nucleic acid sequence encoding a protein to pregulation of expression of a nucleic acid sequence encoding a protein to pregulation of expression of a nucleic A protein kinase is useful for screening biological agents that modulate cancer associated protein kinase function. Downregulating the activity of protein kinase is useful for inhibiting the growth of a cancer call, e.g. liver or colon cancer. A nucleic acid encoding protein kinase is useful to screen biopsy of erived tumours and inflammatory samples such as arthritic synorium, for amplified DNA in the cell or increased expression of corresponding meNA or protein and is also useful to detect differences in expression levels two cells. The present sequence is human mitogen activated protein kinase MAD3XII gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, cytostatic, antisense gene therapy; screening; protein kinase; cancer; liver; colon; tumour; inflammation; arthritic synovium; MAP3K11; mitogen activated protein kinase; enzyme; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diagnosing cancer, comprises determining the upregulation of expression of a nucleic acid sequence encoding a protein kinase or upregulation of expression of the protein kinase, in the cancer
1196 CACCGTGATCTCAAGTCCCAACAACATTTTGCTGCTGCAGCCCATTGAGAGTGACGACATG 1255
                                                                62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                     1316 CAAATGAGTGCCGCGGGCACCTACGCCTGGATGCTCTGAGG 1358
                                                                                                                                                                                                                                                                                                                                                                                                                      Human mitogen activated protein kinase, MAP3K11 gene.
                                                                                                                                122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /product= "Human MAP3K11 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 16; Page 56-60; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                      AAD36139 standard; DNA; 3558 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KINE-) KINETEK PHARM INC.
(UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-0CT-2000; 2000US-237419P.
02-0CT-22000; 2000US-23423P.
04-0CT-2000; 2000US-238558P.
10-MAY-2001; 2001US-290555P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-2001; 2001WO-1B02237.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-233999P.
                                                                                                                                                                                                                                                                                                                                                                            09-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yoganathan T, Delaney AD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-394145/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAE22763.
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02-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAR-2002.
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                                                                                                                                                                                                                                           RESULT 10
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31-JAN-2000, 2000US-0179065.

24-FEB-2000, 2000US-0186528.

25-FEB-2000, 2000US-01865350.

15-FARR-2000, 2000US-01865351.

16-FARR-2000, 2000US-0186351.

19-FARR-2000, 2000US-019874.

11-MAR-2000, 2000US-019874.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218136.

11-MAR-2000, 2000US-0218290.

11-MAR-2000, 2000US-0218290.

11-MAR-2000, 2000US-0218290.

14-MAG-2000, 2000US-0218291.

14-MAG-2000, 2000US-0228281.

15-FEP-2000, 2000US-022891.

16-FEP-2000, 2000US-0228141.

16-FEP-2000, 2000US-0228141.

16-FEP-2000, 2000US-0231243.

16-FEP-2000, 2000US-0231243.

16-FEP-2000, 2000US-0231243.

18-FEP-2000, 2000US-0231243.
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; 2000US-0232398.
; 2000US-0232399.
; 2000US-0232400.
; 2000US-0232401.
                                                                                                                                                                                                                                        17-JAN-2001; 2001WO-US01334
                                                                                                                                                                     WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-SEP-2000;
14-SEP-2000;
14-SEP-2000;
                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                       16-AUG-2001.
The present invention describes a method (M1) for screening for an anti-nepulastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in expression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 844 7 sequences (given in ABLOi664 to ABLO010), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has optostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a structure and/or properties of the agent. M1 can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, ossophageal, ovarian, kidney, prostate or pancraatic cancer, cesophageal, oxarian, acarianna, clear cell cancer, infiltrating ductal cancer, infiltrating ductal cancer, infiltrating ductal cancer, infiltrating lobular cancer, sectionma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1006 GCAGACACGGTGCTCAAGATCACGGACTTCGGCCTCGCCGCGGGGGGCACAAGACCACC 1065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                      Augustus M. Carter KC, Ebner R, Endress G, Horrigan S; Weaver \mathbb{Z}_{2}
                                                                                                                                                                                                                                                                                                                                      Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene set
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Owery Match 44.8%; Score 73.4; DB 24; Length 3454; Best focal Similarity 65.6%; Pred. No. 2.5e-11; Matches 107; Conservative 0; Mismatches 55; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human nervous system related polynucleotide SEQ ID NO 9508.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3454 BP; 594 A; 1217 C; 1136 G; 507 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1066 AAGATGAGCGCTGCGGGGACCTACGCCTGGATGCGCCGGGAGG 1108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID 8355; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA17177 standard; DNA; 7084 BP
                                          02-0CT-2000, 2000US-237295P.
02-0CT-2000, 2000US-237316P.
03-0CT-2000, 2000US-237498P.
03-0CT-2000, 2000US-237694P.
03-0CT-2000, 2000US-237664P.
03-0CT-2000, 2000US-237666P.
01-WOY-2000, 2000US-237666P.
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                                                                                                                                                                                                                        (AVAL-) AVALON PHARM.
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                                                                                                                                                                                                                                                                       Soppet DR,
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Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immuosuppressive; antiinflammatory; anti-HTV; antibacterial; vulnerary; antiparkinsonian; antistching; antianaemic; antiarthritic; cancer; antistching; cerebroprotective; antiinflammatory; antialergic; antidabelic; antiolicer; anticonvisant; antiingal; antialergic; antidabelic; antiologic; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine; ds.
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2000US-0233063.
2000US-0233064.
2000US-0233065.
2000US-0234223.
2000US-0234274.
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2000US-0235484
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2000US-0251856
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                               14 - SEP - 2000; 25 - SEP - 2000; 25 - SEP - 2000; 26 - SEP - 2000; 27 - SEP - 2000; 27 - SEP - 2000; 27 - SEP - 2000; 29 - SEP - 2000; 20 - S
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(ABB14678-ABB18001) useful for grace (introduction) and procession medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune throiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as mycoardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and parasitic infections.
                                                                                                                                                                                                                                                    Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4670 TTTTGCTGCTGCAGCCCATTGAGAGTGACGACATGGAGCACAAGACCCTGAAGATCACCG 4729
                                                                                                                                                                                                                                                                                                                                                                     invention relates to novel genes (ABA11004-ABA21534) and proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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44.1%; Score 72.4; DB 22; Length 7084;
Best Local Similarity 70.3%; Pred. No. 7.3e-14;
Matches 97; Conservative 0; Mismatches 41; Indels 0;
                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 9508; 1701pp + Sequence Listing; English
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                    2000US-0251869.
2000US-0251989.
2000US-0251990.
2000US-0254097.
2001US-0259678.
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  2000US-0251868
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20000S-0250160.
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  20000S - 0.186528
20000S - 0.18664
20000S - 0.18664
20000S - 0.189123
20000S - 0.205515
20000S - 0.205515
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20000S-0231243
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20000S-0231241
20000S-0231414
20000S-0232081
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20000S-0227009.
20000S-0228924.
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2000US-0225759.
2000US-0226279.
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2000US-0235836.
2000US-0236327.
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Ruben SM

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                                       Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4670 TITIGCIGCIGCAGCCCATIGAGAGTGACGACATGGAGCACAAGACCCTGAAGATCACG 4729
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/note= "base labelled as X in the specification"
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/note= "base labelled as X in the specification"
                                                                                                                             Disclosure; SEQ ID NO 28728; 3071pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7084 BP; 1519 A; 1987 C; 2073 G; 1505 T; 0 other;
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WPI; 2001-483426/52
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The LpTKs are expressed in lymphocytic cells, as well as megakaryocytic cells. Sequencing of LpTK-4 revealed the sequences given in AAQ49751 and AAQ49755. The protein sequence corresp. to AAQ49751 is claimed (claim 7) and stated as given in the specification,
                                                                                                                                                                                                                                                                                                                                                                                                     pTK genes were identified using two sets of degenerative oligonucleotide primers: a first set which amplifies all pTK DNA segments (AAQ49743-44), and a second set which amplifies highly conserved sequences present in the catalytic domain of the c-kit subgroup of pTks (AAQ49745-46). The pTK genes identified are described in AAQ49747-57 and AAR41897-02.
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/note= "base n at position 212 is not identified
in the specification"
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/*tag= b
/note= "base n at position 22 is not identified
in the specification"
                                                                                                                                                                                                                                                                         New protein tyrosine kinase genes and proteins encoded by genes are of human mega-karyocytic origin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein tyrosine-kinase; pTK; LpTK4; agonist; cell growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         43.9%; Score 72; DB 14; Length 40.
69.6%; Pred. No. 2.9e-14;
tive 0; Mismatches 42; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 404 BP; 95 A; 106 C; 123 G; 77 T; 3 other;
                                                                                                                                                            Scadden D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein tyrosine-kinase LpTK4 DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  however is missing from the publication.
                                                                                                       (NEWE-) NEW ENGLAND DEACONESS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                               Groopman J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                         Claim 2; Fig 6; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 69.6%;
Conservative
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                                                     92US-0826935
93WO-US00586
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                                                                                                                                                               Cowley S,
                                                                                                                                                                                                                    WPI; 1993-320330/40.
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misc_difference 212
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les 96; Conserv
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  22-JAN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             103 TTTGCTGCTGCAGCCCATTGAGAGAGAGACATGGAGACATAGAGATCACG 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87 ATTITGGGTTGGCGAGGGAATGGCACAGGACCACCAAAATGAGCACAGGAGGCACCTATG 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                            Agonist antibodies which activate specific protein tyrosine kinase(s) - also activate chimeric proteins of kinase extracellular domain and Ig constant domain, useful for studying, and therapeutic modulation of, cell growth and differentiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27 ITTIGCTACTIGAGAAGATAGAACAIGATGACATCIGCAATAAAACTITGAAGATTACAG 86
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in the specification"
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                                                                                                                                                                                                                                Lee JM, Matthews W, Tsai SP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 404 BP; 95 A; 106 C; 123 G; 77 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 58-59; 125pp; English.
                                                                                                                                                                94US-0222616.
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                                                                                                                                                                                                                                Bennett BD, Goeddel D,
                                                                                                                                                                                                (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                               WPI; 1995-366160/47.
                                                                                                                                04-APR-1995;
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